ELSEVIER

Virus Research 38 (1995) 319–321

Virus Research

Author Index

Adams, M.J., see Shi, N.-n. (38) 193
Akkina, R.K., see Sullivan, D.G. (38) 231
Antoniw, J.F., see Shi, N.-n. (38) 193
Apekina, N., see Plyusnin, A. (38) 25

Barge, A., see Di Guilmi, A.M. (38) 71
Bassit, L., see Stuyver, L. (38) 137
Betenbaugh, M., Yu, M., Kuehl, K., White, J., Pennock, D., Spik, K. and Schmaljohn, C.
Nucleocapsid- and virus-like particles assemble in cells infected with recombinant baculoviruses or vaccinia viruses expressing the M and the S segments of Hantaan virus (38) 111
Bifano, M., see Sheaffer, A.K. (38) 305
Bowen, M.D., Kariwa, H., Rollin, P.E., Peters, C.J. and Nichol, S.T.
Genetic characterization of a human isolate of Puumala hantavirus from France (38) 279
Briedis, D.J., see Liston, P. (38) 241
Britton, P., see Chen, C.-M. (38) 83
Brown, B., see Kopecka, H. (38) 125
Brown, W.C., see Li, J. (38) 93
Brummer-Korvenkontio, M., see Plyusnin, A. (38) 25

Carpino, M.R., see Li, J. (38) 93
Casal, J.I., see Laviada, M.D. (38) 205
Cavanagh, D., see Chen, C.-M. (38) 83
Chen, C.-M., Cavanagh, D. and Britton, P.
Cloning and sequencing of a 8.4-kb region from the 3′-end of a Taiwanese virulent isolate of the coronavirus transmissible gastroenteritis virus (38) 83
Chen, J., see Shi, N.-n. (38) 193
Chroboczek, J., see Di Guilmi, A.M. (38) 71
Collisson, E.W., see Li, J. (38) 93
Colonno, R.J., see Sheaffer, A.K. (38) 305

DiFlumeri, C., see Liston, P. (38) 241
Di Guilmi, A.M., Barge, A., Kitts, P., Gout, E. and Chroboczek, J.
Human adenovirus serotype 3 (Ad3) and the Ad3 fiber protein bind to a 130-kDa membrane protein on HeLa cells (38) 71
Doerfler, W., see Orend, G. (38) 261

Elder, J.H., see Li, J. (38) 93

Foulds, I.J., see Shi, N.-n. (38) 193

Gavrilovskaya, I., see Plyusnin, A. (38) 25
Gout, E., see Di Guilmi, A.M. (38) 71
Grant, C.K., see Li, J. (38) 93
Gray, W.L., see Pumphrey, C.Y. (38) 55

Hamatake, R.K., see Sheaffer, A.K. (38) 305
Han, M.H., see Jin, B.R. (38) 269
Henttonen, H., see Plyusnin, A. (38) 25
Hong, H.J., see Jin, B.R. (38) 269
Hongo, S., see Matsuzaki, Y. (38) 291
Hosaka, Y., see Kuroda, K. (38) 43
Hurlburt, W.W., see Sheaffer, A.K. (38) 305

Ikeda, T., Takase-Yoden, S. and Watanabe, R.
Characterization of monoclonal antibodies recognizing neurotropic Friend murine leukemia virus (38) 297

Jang, H.-K., see Ono, M. (38) 219
Jin, B.R., Ryu, C.J., Kang, S.K., Han, M.H. and Hong, H.J.
Characterization of a murine–human chimeric antibody with specificity for the pre-S2 surface antigen of hepatitis B virus expressed in baculovirus-infected insect cells (38) 269

Kang, S.K., see Jin, B.R. (38) 269
Kariwa, H., see Bowen, M.D. (38) 279
Kawaguchi, Y., see Ono, M. (38) 219
Kitame, F., see Matsuzaki, Y. (38) 291
Kitts, P., see Di Guilmi, A.M. (38) 71
Kleter, B., see Stuyver, L. (38) 137
Knoblauch, M., see Orend, G. (38) 261
Kopecka, H., Brown, B. and Pallansch, M.
Genotypic variation in Coxsackievirus B5 isolates from three different outbreaks in the United States (38) 125
Kuehl, K., see Betenbaugh, M. (38) 111
Kuroda, K., Nakajima, M., Minakata, H. and Hosaka, Y.
Identification of K^k-restricted T-cell epitope within influenza virus nucleoprotein (38) 43

Laakkonen, J., see Plyusnin, A. (38) 25
Laurent-Puig, P., see Stuyver, L. (38) 137
Laviada, M.D., Roy, P., Sánchez-Vizcaíno, J.M. and Casal, J.I.
The use of African horse sickness virus NS3 protein, expressed in bacteria, as a marker to differentiate infected from vaccinated horses (38) 205
Lehväslaiho, H., see Plyusnin, A. (38) 25
Li, J., Brown, W.C., Song, W., Carpino, M.R., Wolf, A.M., Grant, C.K., Elder, J.H. and Collisson, E.W.
Retroviral vector-transduced cells expressing the core polyprotein induce feline immunodeficiency virus-specific cytotoxic T-lymphocytes from infected cats (38) 93
Liston, P., DiFlumeri, C. and Briedis, D.J.
Protein interactions entered into by the measles virus P, V, and C proteins (38) 241
Lunel, F., see Stuyver, L. (38) 137

MacFarlane, S.A., see Shi, N.-n. (38) 193
Maeda, K., see Ono, M. (38) 219
Maertens, G., see Stuyver, L. (38) 137
Mateu, M.G.
Antibody recognition of picornaviruses and escape from neutralization: a structural view (38) 1
Matsuzaki, M., see Matsuzaki, Y. (38) 291
Matsuzaki, Y., Matsuzaki, M., Muraki, Y., Sugawara, K., Hongo, S., Kitame, F. and Nakamura, K.
Comparison of receptor-binding properties among influenza C virus isolates (38) 291
Mikami, T., see Ono, M. (38) 219
Mikhailova, T., see Plyusnin, A. (38) 25
Minakata, H., see Kuroda, K. (38) 43
Morzunov, S.P., Winton, J.R. and Nichol, S.T.
The complete genome structure and phylogenetic relationship of infectious hematopoietic necrosis virus (38) 175
Muraki, Y., see Matsuzaki, Y. (38) 291

Nakajima, M., see Kuroda, K. (38) 43
Nakamura, K., see Matsuzaki, Y. (38) 291
Nichol, S.T., Rowe, J.E. and Winton, J.R.
Molecular epizootiology and evolution of the glycoprotein and non-virion protein genes of infectious hematopoietic necrosis virus, a fish rhabdovirus (38) 159
Nichol, S.T., see Bowen, M.D. (38) 279
Nichol, S.T., see Morzunov, S.P. (38) 175
Niemimaa, J., see Plyusnin, A. (38) 25
Niikura, M., see Ono, M. (38) 219
Nkengasong, J., see Stuyver, L. (38) 137

Ono, M., Jang, H.-K., Maeda, K., Kawaguchi, Y., Tohya, Y., Niikura, M. and Mikami, T.
Preparation of monoclonal antibodies against Marek's disease virus serotype 1 glycoprotein D expressed by a recombinant baculovirus (38) 219
Orend, G., Knoblauch, M. and Doerfler, W.
Selective loss of unmethylated segments of integrated Ad12 genomes in revertants of the adenovirus type 12-transformed cell line T637 (38) 261

Pallansch, M., see Kopecka, H. (38) 125
Pawlotsky, J.-M., see Stuyver, L. (38) 137
Pennock, D., see Betenbaugh, M. (38) 111
Peters, C.J., see Bowen, M.D. (38) 279
Plyusnin, A., Vapalahti, O., Lehväslaiho, H., Apekina, N., Mikhailova, T., Gavrilovskaya, I., Laakkonen, J., Niemimaa, J., Henttonen, H., Brummer-Korvenkontio, M. and Vaheri, A.
Genetic variation of wild Puumala viruses within the serotype, local rodent populations and individual animal (38) 25
Pumphrey, C.Y. and Gray, W.L.
DNA sequence of the simian varicella virus (SVV) gH gene and analysis of the SVV and varicella zoster virus gH transcripts (38) 55

Rollin, P.E., see Bowen, M.D. (38) 279
Rowe, J.E., see Nichol, S.T. (38) 159
Roy, P., see Laviada, M.D. (38) 205
Ryu, C.J., see Jin, B.R. (38) 269

Sánchez-Vizcaíno, J.M., see Laviada, M.D. (38) 205

Schmaljohn, C., see Betenbaugh, M. (38) 111
Sheaffer, A.K., Hurlburt, W.W., Stevens, J.T., Bifano, M., Hamatake, R.K., Colonno, R.J. and Tenney, D.J.
Characterization of monoclonal antibodies recognizing amino- and carboxy-terminal epitopes of the herpes simplex virus UL42 protein (38) 305
Shi, N.-n., Zhu, M., Chen, J., Stratford, R., Wilson, T.M.A., Antoniw, J.F., Foulds, I.J., MacFarlane, S.A. and Adams, M.J.
Molecular characterisation of UK isolates of barley yellow mosaic bymovirus (38) 193
Song, W., see Li, J. (38) 93
Spik, K., see Betenbaugh, M. (38) 111
Stevens, J.T., see Sheaffer, A.K. (38) 305
Stratford, R., see Shi, N.-n. (38) 193
Stuyver, L., Wyseur, A., Van Arnhem, W., Lunel, F., Laurent-Puig, P., Pawlotsky, J.-M., Kleter, B., Bassit, L., Nkengasong, J., Van Doorn, L.-J. and Maertens, G.
Hepatitis C virus genotyping by means of 5′-UR/core line probe assays and molecular analysis of untypeable samples (38) 137
Sugawara, K., see Matsuzaki, Y. (38) 291
Sullivan, D.G. and Akkina, R.K.
A nested polymerase chain reaction assay to differentiate pestiviruses (38) 231

Takase-Yoden, S., see Ikeda, T. (38) 297
Tenney, D.J., see Sheaffer, A.K. (38) 305
Tohya, Y., see Ono, M. (38) 219

Vaheri, A., see Plyusnin, A. (38) 25
Van Arnhem, W., see Stuyver, L. (38) 137
Van Doorn, L.-J., see Stuyver, L. (38) 137
Vapalahti, O., see Plyusnin, A. (38) 25

Watanabe, R., see Ikeda, T. (38) 297
White, J., see Betenbaugh, M. (38) 111
Wilson, T.M.A., see Shi, N.-n. (38) 193
Winton, J.R., see Morzunov, S.P. (38) 175
Winton, J.R., see Nichol, S.T. (38) 159
Wolf, A.M., see Li, J. (38) 93
Wyseur, A., see Stuyver, L. (38) 137

Yu, M., see Betenbaugh, M. (38) 111

Zhu, M., see Shi, N.-n. (38) 193

ELSEVIER

Virus Research 38 (1995) 323–324

Virus Research

Subject Index

Adenovirus, (38) 71
Adenovirus oncogenesis, (38) 261
African horse sickness virus, (38) 205
Antibody recognition, (38) 1

Baculovirus, (38) 111
Baculovirus expression system, (38) 71
Baculovirus recombinant, (38) 219
Baculovirus vector, (38) 269
Barley yellow mosaic virus, (38) 193
B5-genotypic variation, (38) 125
Border disease virus, (38) 231
Bovine viral diarrhea virus, (38) 231
Bunyaviridae, (38) 279

Cellular receptor, (38) 71
Chimeric antibody, (38) 269
Conformation-dependent recognition, (38) 297
Core polyprotein, (38) 93
Coronavirus, (38) 83
Coxsackievirus, (38) 125
C protein, (38) 241
Cytotoxic T-lymphocyte, (38) 43, (38) 93

Differential ELISA, (38) 205
DNA methylation, (38) 261
DNA replication, (38) 305

Escape mutation, (38) 1
Evolution, (38) 25
Expression, (38) 269

Feline immunodeficiency virus (FIV), (38) 93
Fiber protein, (38) 71
Fish disease, (38) 159
Fish rhabdovirus, (38) 159, (38) 175
Foot-and-mouth disease virus, (38) 1
Friend murine leukemia virus, (38) 297

gag, (38) 93
Genetic variation, (38) 25
Glycoprotein D, (38) 219
Glycoprotein H gene, (38) 55

Hantaan virus, (38) 111
Hantavirus, (38) 25, (38) 279
HCV genotyping, (38) 137
Hemagglutinin-esterase, (38) 291
Hepatitis B virus, (38) 269
Herpes simplex virus, (38) 305
Host cell protein, (38) 241

Immune response, (38) 55
Infectious hematopoietic necrosis virus, (38) 159, (38) 175
Influenza virus, (38) 43, (38) 291
Integrated viral genome, (38) 261
Isolate difference, (38) 193

Line probe assay, (38) 137
Loss of integrated viral genome, (38) 261
L polymerase, (38) 175

Marek's disease virus serotype 1, (38) 219
Measles virus, (38) 241
Membrane protein, (38) 71
Monoclonal antibody, (38) 219, (38) 297, (38) 305

Nested polymerase chain reaction (PCR), (38) 231
Neurotropism, (38) 297
NS3 protein, (38) 205

Peripheral blood mononuclear cell (PBMC), (38) 93
Pestiviral genotype, (38) 231

Pestivirus, (38) 231
Phosphoprotein (P protein), (38) 241
Phylogenetic analysis, (38) 137, (38) 175
Picornavirus, (38) 1
Porcine, (38) 83
Protein interaction, (38) 241
Puumala virus, (38) 25, (38) 279

Quasispecies, (38) 25

Receptor-binding site, (38) 291
Resistance-breaking strain, (38) 193
Retroviral vector, (38) 93
Rhabdoviridae, (38) 175
RNA, medium, (38) 279
RNA sequence, (38) 175
RNA, small, (38) 279

Simian varicella virus, (38) 55
Swine, (38) 83

T-cell epitope, (38) 43
T-lymphocyte, (38) 93
Transmissible gastroenteritis virus (TGEV), (38) 83

Vaccinia virus, (38) 111
Varicella zoster virus, (38) 55
Virus evolution, (38) 159
Virus-like particle, (38) 111
V protein, (38) 241